18

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PCT/FR00/01887

SEQUENCE LISTING

(1105 INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDIC. [NATIONAL INSTITUTE FOR HEALTH AND MEDICAL RESEARCH] - INSERM	
<pre><120 + D18210</pre>	
$0.030 + 0.08$ of the $\mathit{Krit1}$ gene in the field of angiogenesis	
0140 + 0141 +	
<12 kg D + 15 11	
<170 - PatentIn Vers. 2.0	
<pre><210 > 1 <211 + 18 <211 + ENA <213 + Artificial Sequence</pre>	
<223 - Jense primer	
<pre>c400 - 1 gagaggataa aaactaat</pre>	18
<pre><210 - 3 <211 - 18 <!--112 - DNA <2113 - Artificial Sequence</pre--></pre>	
<pre><000 + <000 + <000 + Reverse primer</pre>	
<400 · C ywystawaat toattoaa	18
<pre><:10 - 3 <:111 - 18 <:111 - DNA</pre>	

€2200 ×

Sense primer

44000 €

gotottaatg ggtttttg

+1.11.55.4

0.000 4
6.0010 18
6.0010 ENA
6.0130 Artificial Sequence

<:::: Artificial Sequence</pre>

 $< 0.00 \leq 0.00 \leq$

<1223> Feverse primer

18

*: Artificial Sequence

HERE Sense primer

atgtaatgcc tttttcc

1200

<4000-9

<2100 10 <2110 18

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:212 - DNA ::213 - Artificial Sequence		
+(2.20 + +(2.23 + Reverse primer		
(40) + 10 atjustggot otaactat		18
HC10 + 11 HC11 + 13 HC13 + DNA HC13 + Artificial Sequence		
+1/2005 +1223 - Sense primer		
<400 × 11 tiyttagatt gtgatgta		18
HD16 - 12 HD11 - 10 HD12 - ENA HD13 - Artificial Sequence		
<pre><310 > <3213 > Feverse primer</pre>		
<pre><i400 10="" aaactttc<="" ascataataa="" pre="" ×=""></i400></pre>		18
<pre><c108 13="" 18="" <c118="" <c138="" artificial="" ena="" pre="" sequence<=""></c108></pre>		
<pre><pre><pre></pre> <pre></pre> <pre>Sense primer</pre></pre></pre>		
<pre><400 > 13 titalaaaag gaatgatg</pre>		18
<pre><cli>+Cli0 + 14 <cli>+Cli + 18 <cli>+Cli3 > DNA <cli>+Cli3 > Artificial Sequence</cli></cli></cli></cli></pre>		
<pre></pre> <pre><pre></pre> <pre></pre> <pre></pre> <pre></pre> <pre>Reverse primer</pre></pre>		
-:400 - 14 toaactcaaa ccatatca		18

+1110 - 15 +111 + 18 +1211 + DNA +1218 - Artificial Sequence

<0000> <0000>

10 01/02004	101/1800/0188
<pre>44) 15 tgtw:cotaa taaccaaa</pre>	18
+(21) + 16 +(21) + 18 +(21) + 1MA +(31) + Artificial Sequence	
<pre>*Min * *Min * Peverse primer</pre>	
रक्षण र 16 अनुदर्भ ay sac aagaccat	18
Hill 17 Hill: 18 Hill: DNA Hill: Artificial Sequence	
HUCO - Hanse primer	
8400×17 gatawagitt itaatatg	18
#210 + 18 #211 + 18 #211 + EMA #213 + Artificial Sequence	
<pre><!--D0 + <!D0 + Feverse primer</pre--></pre>	
<400 - 18 caatagttta tgaagtoo	18
<pre><c10> 19 <c11 +="" 18="" <c11="" <c13="" artificial="" pma="" pre="" sequence<=""></c11></c10></pre>	
HIDE + Sonse primer	
<pre>c400 - 10 atatttasaa aggcaage</pre>	18
<pre><210 + 20 <2210 + 18 <2210 + DMA <2213 - Artificial Sequence</pre>	
KULC KULC - Feverse primer	
k400+30 tqacatqatt ggtaaaaa	18
<pre><pre><pre><pre><pre>< 21</pre></pre></pre></pre></pre>	

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PRIOR MA	
- 1: Antificial Sequence	
•	
- 224 c	
: :: :: :: :: :: :: :: : :: : :: :	
9.400 × 21	
tygtadeitt tootttoa	18
+210 + 23	
1.111 · 14	
- C1 INA	
+::: Artificial Sequence	
-L.F. Feverse primer	
k490% 22	
etttatgatt getgggge	18
+2009-23	
+ 111 x 16	
+ 212 + CHA	
-213 - Artificial Sequence	
0223% Sense primer	
+4000 (33	
	18
gatgaagstt ttaatatg	10
K210% 24	
×211× 19	
+ 212 + CNA	
1139 Artificial Sequence	
211 Mollitoral Stydenes	
+ 32 G/A	
<pre>#D13# Reverse primer</pre>	
• -	
+4602-14	
meatagtitta tgaagtoo	18
+ 210 + 15	
V2113-18	
+ 010 + ENA	
+113+ Artificial Sequence	
H220 K	
#123 - Sense primer	
ALCO OF	
- 1400 + 25 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 -	1.0
aalajatagg gaactgoo	18
Fig. 10 - 26	
+111 + 16	
· III. PNA	
· 211 · Artificial Sequence	
Y. 10 A	
+213 · Feverse primer	

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-213 - Homo sapiens		
-40) - 35 Taatmattag ggaacgacag		20
- 0100-34 - 0110-00 - 0100-00A		
-21 Hemo sapiens		
-4005 k4 anghatgoty gtaaatggaa		20
- 210 - 35 - 210 - 20 - 210 - 20A		
-2130 Home sapiens		
-4002-35 ttttatabag gtatggaaaa		<u>:</u> 0
+ 210 + 36 + 211 + 20 + 210 + BDA		
- 213: Homo sapiens		
-400 - 36 Sacqgataga gtaagttatt		20
-2100-37 -211>-20 -210>-DNA -213>-Homo sapiens		
±400≥ 37		
Abanticiag catataacag + 2109 38		.:0
+2115 20 +2135 DNA		
- 213 - Homo sapiens - 400- 3-		
taacaasca gtaagaatta		0
+ 210% ±0 + 211% 20 + 21.% DNA		
-213: Hamo sapiens		
-400 -34 tutontgrag tatgaaaaag		20
+ 210 + 41 + 211 + 20		
+212 + FMA +213 + Hamo sapiens		

20

(400) 40 gaadacctca gtaagaaagt -:.100 - 41

-.. 11:- 2:0 -:::1:::- DNA

Hul Homo sapiens

-14001-41

tottitt dag goottoaact

*1.1100 - 4.70

-0110- 10 -0110- DNA

-1111 Homo sapiens

-140m. - 4.1

igaaaaarag gtttgcttgg

HII 100 4 3

·:211: 20

-121.1 - LNA

Mills Himo sapiens

-14000-43

inocittaad attgaagacc

SIL101-44

*2110 20 *2110 EWA

+:2150 Hamo sapiens

-4005 44

atticciaaa gtaagtattt

+12.100 + 4.5

H211- 20

HILLS DUA

+1113 - Homo sapiens

44000 45

sigettaeag tgaagaaaat

H1010 - 46

+12111- 00

- 11. ENA

+1213 - Homo sapiens

*:400.46

tqaatacaag gtaagctgtt

+1110 + 47

P211-10

-0.112 - FNA

>0.113 * Home sapiens

4499 47

rigittitaq aatotoagta

 $\pm 1210 \pm 48$

<211 - 20

HIIII HIMA

+:213. Homo sapiens

20

20

20

20

20

20

20

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tactttgtag gctctggtcg